

EXHIBIT A PENDING CLAIMS (SERIAL NO. 08/951,188; 4200.000200)

- 110. (Amended) An isolated nucleic acid molecule comprising a nucleic acid sequence that encodes:
 - (a) a polypeptide having the amino acid sequence of SEQ ID NO:2; or
 - (b) a polypeptide that comprises a contiguous sequence of at least 16 amino acids from SEQ ID NO:4, of at least 20 amino acids from SEQ ID NO:45, of at least 20 amino acids from SEQ ID NO:47 or of at least 125 amino acids from SEQ ID NO:50.
- 111. The isolated nucleic acid molecule of claim 110, comprising a nucleic acid sequence that encodes a polypeptide having the amino acid sequence of SEQ ID NO:2.
- 112. The isolated nucleic acid molecule of claim 111, comprising a nucleic acid sequence that has the nucleotide sequence from position 115 to position 1327 of SEQ ID NO:1.
- 113. (Amended) An isolated nucleic acid molecule comprising a nucleic acid sequence that encodes a polypeptide that comprises a contiguous sequence of at least 16 amino acids from SEQ ID NO:4, of at least 20 amino acids from SEQ ID NO:45, of at least 20 amino acids from SEQ ID NO:47 or of at least 125 amino acids from SEQ ID NO:50.
- 116. (Amended) The isolated nucleic acid molecule of claim 113, comprising a nucleic acid sequence that encodes a polypeptide that comprises a contiguous sequence of at least 25 amino acids from SEQ ID NO:4, SEQ ID NO:45 or SEQ ID NO:47.
- 117. (Amended) The isolated nucleic acid molecule of claim 116, comprising a nucleic acid sequence that encodes a polypeptide that comprises a contiguous sequence of at least about 30 amino acids from SEQ ID NO:4, SEQ ID NO:45 or SEQ ID NO:47.
- 118. (Amended) The isolated nucleic acid molecule of claim 117, comprising a nucleic acid sequence that encodes a polypeptide that comprises a contiguous sequence of at least about 40 amino acids from SEQ ID NO:4, SEQ ID NO:45 or SEQ ID NO:47.

- 119. (Amended) The isolated nucleic acid molecule of claim 118, comprising a nucleic acid sequence that encodes a polypeptide that comprises a contiguous sequence of at least about 50 amino acids from SEQ ID NO:4, SEQ ID NO:45 or SEQ ID NO:47.
- 120. (Amended) The isolated nucleic acid molecule of claim 119, comprising a nucleic acid sequence that encodes a polypeptide that comprises a contiguous sequence of at least about 60 amino acids from SEQ ID NO:4, SEQ ID NO:45 or SEQ ID NO:47.
- 121. (Amended) The isolated nucleic acid molecule of claim 120, comprising a nucleic acid sequence that encodes a polypeptide that comprises a contiguous sequence of at least about 70 amino acids from SEO ID NO:4, SEO ID NO:45 or SEO ID NO:47.
- 122. (Amended) The isolated nucleic acid molecule of claim 121, comprising a nucleic acid sequence that encodes a polypeptide that comprises a contiguous sequence of at least about 80 amino acids from SEQ ID NO:4, SEQ ID NO:45 or SEQ ID NO:47.
- 123. (Amended) The isolated nucleic acid molecule of claim 122, comprising a nucleic acid sequence that encodes a polypeptide that comprises a contiguous sequence of at least about 90 amino acids from SEQ ID NO:4, SEQ ID NO:45 or SEQ ID NO:47.
- 124. (Amended) The isolated nucleic acid molecule of claim 123, comprising a nucleic acid sequence that encodes a polypeptide that comprises a contiguous sequence of at least about 100 amino acids from SEQ ID NO:4, SEQ ID NO:45 or SEQ ID NO:47.
- 125. (Amended) The isolated nucleic acid molecule of claim 124, comprising a nucleic acid sequence that encodes a polypeptide that comprises a contiguous sequence of at least 125 amino acids from SEQ ID NO:4, SEQ ID NO:45, SEQ ID NO:47 or SEQ ID NO:50.
- 126. The isolated nucleic acid molecule of claim 125, comprising a nucleic acid sequence that encodes a polypeptide that comprises a contiguous sequence of at least about 150 amino acids from SEQ ID NO:4, SEQ ID NO:45, SEQ ID NO:47 or SEQ ID NO:50.
- 127. The isolated nucleic acid molecule of claim 126, comprising a nucleic acid sequence that encodes a polypeptide that comprises a contiguous sequence of at least about 200 amino acids from SEQ ID NO:4, SEQ ID NO:45, SEQ ID NO:47 or SEQ ID NO:50.

- 128. The isolated nucleic acid molecule of claim 127, comprising a nucleic acid sequence that encodes a polypeptide that comprises the amino acid sequence of SEQ ID NO:4, SEQ ID NO:45, SEQ ID NO:47 or SEQ ID NO:50.
- 129. The isolated nucleic acid molecule of claim 128, comprising a nucleic acid sequence that encodes a polypeptide that comprises the amino acid sequence of SEQ ID NO:4.
- 130. The isolated nucleic acid molecule of claim 128, comprising a nucleic acid sequence that encodes a polypeptide that comprises the amino acid sequence of SEQ ID NO:45.
- 131. The isolated nucleic acid molecule of claim 128, comprising a nucleic acid sequence that encodes a polypeptide that comprises the amino acid sequence of SEQ ID NO:47.
- 132. The isolated nucleic acid molecule of claim 128, comprising a nucleic acid sequence that encodes a polypeptide that comprises the amino acid sequence of SEQ ID NO:50.
- 133. The isolated nucleic acid molecule of claim 113, wherein said nucleic acid sequence is operatively positioned under the control of a promoter.
- 134. The isolated nucleic acid molecule of claim 133, further defined as a recombinant vector.
- 135. The isolated nucleic acid molecule of claim 133, comprised within a recombinant host cell.
- 136. The isolated nucleic acid molecule of claim 113, wherein said nucleic acid sequence is operatively attached to a second coding region that encodes a selected peptide or protein sequence so that said isolated nucleic acid molecule encodes a fusion protein.
- 137. (Amended) An isolated nucleic acid molecule comprising a nucleic acid sequence that encodes a P-TEFb large subunit protein, wherein said P-TEFb large subunit protein binds to a P-TEFb kinase subunit protein to form a P-TEFb enzyme complex that promotes transcription elongation and wherein said nucleic acid molecule comprises the nucleotide sequence of:

the coding sequence of a cDNA molecule present in a nucleic acid library, wherein the cDNA molecule hybridizes to a probe having the sequence of the complement of SEQ ID NO:3, SEQ ID NO:43 or SEQ ID NO:48 under

conditions of high stringency comprising hybridization in 50% formamide, $5 \times$ Denhardts' solution, $5 \times$ SSC, 25 mM sodium phosphate, 0.1% SDS and 100 µg/ml of denatured salmon sperm DNA at 42°C for 16 h followed by 1h sequential washes with 0.1× SSC, 0.1% SDS solution at 60°C.

- 138. The isolated nucleic acid molecule of claim 137, wherein the nucleic acid molecule comprises a nucleotide sequence of at least 21 contiguous nucleotides present in SEQ ID NO:3, SEQ ID NO:43 or SEQ ID NO:48.
- 139. The isolated nucleic acid molecule of claim 138, wherein the nucleic acid molecule comprises a nucleotide sequence of at least 30 contiguous nucleotides present in SEQ ID NO:3, SEQ ID NO:43 or SEQ ID NO:48.
- 140. The isolated nucleic acid molecule of claim 139, wherein the nucleic acid molecule comprises a nucleotide sequence of at least 40 contiguous nucleotides present in SEQ ID NO:3, SEQ ID NO:43 or SEQ ID NO:48.
- 141. The isolated nucleic acid molecule of claim 140, wherein the nucleic acid molecule comprises a nucleotide sequence of at least 50 contiguous nucleotides present in SEQ ID NO:3, SEQ ID NO:43 or SEQ ID NO:48.
- 142. The isolated nucleic acid molecule of claim 141, wherein the nucleic acid molecule comprises a nucleotide sequence of at least 60 contiguous nucleotides present in SEQ ID NO:3, SEQ ID NO:43 or SEQ ID NO:48.
- 143. The isolated nucleic acid molecule of claim 142, wherein the nucleic acid molecule comprises a nucleotide sequence of at least 72 contiguous nucleotides present in SEQ ID NO:3, SEO ID NO:43 or SEO ID NO:48.
- 144. The isolated nucleic acid molecule of claim 137, wherein the nucleic acid molecule has the nucleotide sequence of SEQ ID NO:44.
- 145. The isolated nucleic acid molecule of claim 137, wherein the nucleic acid molecule has the nucleotide sequence of SEQ ID NO:46.

- 146. The isolated nucleic acid molecule of claim 137, wherein the nucleic acid molecule has the nucleotide sequence of SEQ ID NO:49.
- 147. The isolated nucleic acid molecule of claim 137, wherein the nucleic acid molecule is up to about 10,000 basepairs in length.
- 148. The isolated nucleic acid molecule of claim 147, wherein the nucleic acid molecule is up to about 5,000 basepairs in length.
- 149. An isolated nucleic acid molecule comprising a nucleic acid sequence that encodes a P-TEFb large subunit protein that exhibits at least 90% identity to the amino acid sequence set forth in SEQ ID NO:4, SEQ ID NO:45, SEQ ID NO:47 or SEQ ID NO:50, wherein said P-TEFb large subunit protein binds to a P-TEFb kinase subunit protein to form a P-TEFb enzyme complex that promotes transcription elongation.
- 150. The isolated nucleic acid molecule of claim 149, wherein the encoded polypeptide exhibits between 91% and about 95% identity to the amino acid sequence set forth in SEQ ID NO:4, SEQ ID NO:45, SEQ ID NO:47 or SEQ ID NO:50.
- 151. The isolated nucleic acid molecule of claim 150, wherein the encoded polypeptide exhibits between 96% and about 99% identity to the amino acid sequence set forth in SEQ ID NO:4, SEQ ID NO:45, SEQ ID NO:47 or SEQ ID NO:50.
- 152. (Amended) An isolated nucleic acid molecule comprising:
 - (a) a first nucleic acid sequence that encodes a P-TEFb small subunit protein that has kinase activity and binds to a P-TEFb large subunit protein to form a P-TEFb enzyme complex that promotes transcription elongation, wherein said P-TEFb small subunit protein exhibits at least 90% identity to the amino acid sequence set forth in SEQ ID NO:2 or SEQ ID NO:6; and
 - (b) a second nucleic acid sequence that encodes a P-TEFb large subunit protein that binds to a P-TEFb kinase subunit protein to form a P-TEFb enzyme complex that promotes transcription elongation, wherein said P-TEFb large subunit protein exhibits at least 90% identity to the amino acid sequence set forth in SEQ ID NO:4, SEQ ID NO:45, SEQ ID NO:47 or SEQ ID NO:50.

- 153. The isolated nucleic acid molecule of claim 152, wherein said first nucleic acid sequence encodes a polypeptide having the amino acid sequence of SEQ ID NO:6.
- 154. The isolated nucleic acid molecule of claim 152, wherein said second nucleic acid sequence encodes a polypeptide that has the amino acid sequence of SEQ ID NO:45, SEQ ID NO:47 or SEQ ID NO:50.
- 155. The isolated nucleic acid molecule of claim 154, wherein said second nucleic acid sequence has the nucleotide sequence of SEQ ID NO:44, SEQ ID NO:46 or SEQ ID NO:49.
- 156. The isolated nucleic acid molecule of claim 152, wherein said first nucleic acid sequence has the nucleotide sequence of SEQ ID NO:5 and wherein said second nucleic acid sequence has the nucleotide sequence of SEQ ID NO:44, SEQ ID NO:46 or SEQ ID NO:49.
- 157. (Amended) An expression system comprising:
 - (a) a first expression unit comprising, under the transcriptional control of a promoter, a first nucleic acid sequence that encodes a P-TEFb small subunit protein that has kinase activity and binds to a P-TEFb large subunit protein to form a P-TEFb enzyme complex that promotes transcription elongation, wherein said P-TEFb small subunit protein exhibits at least 90% identity to the amino acid sequence set forth in SEQ ID NO:2 or SEQ ID NO:6; and
 - (b) a second expression unit comprising, under the transcriptional control of a promoter, a second nucleic acid sequence as defined in claim 137 or claim 149.
- 158. The expression system of claim 157, wherein said first expression unit comprises a first nucleic acid sequence that encodes a polypeptide that comprises a contiguous sequence of at least about 18 amino acids from SEQ ID NO:2 or SEQ ID NO:6.
- 159. The expression system of claim 158, wherein said first expression unit comprises a first nucleic acid sequence that encodes a polypeptide that comprises a contiguous sequence of at least about 20 amino acids from.
- 160. The expression system of claim 159, wherein said first expression unit comprises a first nucleic acid sequence that encodes a polypeptide that comprises a contiguous sequence of at least about 50 amino acids from SEQ ID NO:2 or SEQ ID NO:6.

- 161. The expression system of claim 160, wherein said first expression unit comprises a first nucleic acid sequence that encodes a polypeptide that comprises a contiguous sequence of at least about 100 amino acids from SEQ ID NO:2 or SEQ ID NO:6.
- 162. The expression system of claim 161, wherein said first expression unit comprises a first nucleic acid sequence that encodes a polypeptide that has the amino acid sequence of SEQ ID NO:2.
- 163. The expression system of claim 161, wherein said first expression unit comprises a first nucleic acid sequence that encodes a polypeptide that has the amino acid sequence of SEQ ID NO:6.
- 164. (Amended) The expression system of claim 157, wherein said second expression unit comprises a second nucleic acid sequence that encodes a polypeptide that comprises a contiguous sequence of at least 16 amino acids from SEQ ID NO:4, of at least 20 amino acids from SEQ ID NO:45, of at least 20 amino acids from SEQ ID NO:47 or of at least 125 amino acids from SEQ ID NO:50.
- 167. (Amended) The expression system of claim 166, wherein said second expression unit comprises a second nucleic acid sequence that encodes a polypeptide that comprises a contiguous sequence of at least about 50 amino acids from SEQ ID NO:4, SEQ ID NO:45 or SEQ ID NO:47.
- 168. (Amended) The expression system of claim 167, wherein said second expression unit comprises a second nucleic acid sequence that encodes a polypeptide that comprises a contiguous sequence of at least about 100 amino acids from SEQ ID NO:4, SEQ ID NO:45 or SEQ ID NO:47.
- 169. The expression system of claim 168, wherein said second expression unit comprises a second nucleic acid sequence that encodes a polypeptide that has the amino acid sequence of SEQ ID NO:4.
- 170. The expression system of claim 168, wherein said second expression unit comprises a second nucleic acid sequence that encodes a polypeptide that has the amino acid sequence of SEQ ID NO:45.

- 171. The expression system of claim 168, wherein said second expression unit comprises a second nucleic acid sequence that encodes a polypeptide that has the amino acid sequence of SEO ID NO:47.
- 172. The expression system of claim 168, wherein said second expression unit comprises a second nucleic acid sequence that encodes a polypeptide that has the amino acid sequence of SEQ ID NO:50.
- 174. (Amended) The expression system of claim 168, wherein said second expression unit comprises a second nucleic acid sequence that has the nucleotide sequence of SEQ ID NO:44.
- 175. (Amended) The expression system of claim 168, wherein said second expression unit comprises a second nucleic acid sequence that has the nucleotide sequence of SEQ ID NO:46.
- 176. (Amended) The expression system of claim 168, wherein said second expression unit comprises a second nucleic acid sequence that has the nucleotide sequence of SEQ ID NO:49.
- 177. (Amended) The expression system of claim 157, wherein said second expression unit comprises a second nucleic acid sequence that encodes a polypeptide that exhibits between 91% and about 95% identity to the amino acid sequence set forth in SEQ ID NO:4, SEQ ID NO:45, SEQ ID NO:47 or SEQ ID NO:50.
- 178. The expression system of claim 157, wherein said first and said second expression units are comprised in a single expression vector.
- 179. The expression system of claim 157, wherein said first and said second expression units are each comprised in a separate expression vector.
- 180. The expression system of claim 157, wherein said expression system is comprised within a recombinant host cell.

- 181. (Amended) An expression system comprising:
 - (a) a first expression unit comprising, under the transcriptional control of a promoter, a first nucleic acid sequence that encodes a polypeptide and that comprises the nucleotide sequence of the coding sequence of a cDNA molecule present in a nucleic acid library, wherein the cDNA molecule hybridizes to a probe having the sequence of the complement of SEQ ID NO:1 or SEQ ID NO:5 under conditions of high stringency comprising hybridization in 50% formamide, 5× Denhardts' solution, 5× SSC, 25 mM sodium phosphate, 0.1% SDS and 100 μg/ml of denatured salmon sperm DNA at 42°C for 16 h followed by 1h sequential washes with 0.1× SSC, 0.1% SDS solution at 60°C; and
 - (b) a second expression unit comprising, under the transcriptional control of a promoter, a second nucleic acid sequence as defined in claim 137 or claim 149.
- 182. (Amended) The expression system of claim 181, wherein said first expression unit comprises a first nucleic acid sequence that comprises the nucleotide sequence of the coding sequence of a cDNA molecule present in a nucleic acid library, wherein the cDNA molecule hybridizes to a probe having the sequence of the complement of SEQ ID NO:5 under said conditions of high stringency.
- 183. (Amended) The expression system of claim 181, wherein said first expression unit comprises a first nucleic acid sequence that comprises the nucleotide sequence of the coding sequence of a cDNA molecule present in a nucleic acid library, wherein the cDNA molecule hybridizes to a probe having the sequence of the complement of the nucleotide sequence from position 115 to position 1327 of SEQ ID NO:1 under said conditions of high stringency.
- 184. The expression system of claim 181, wherein said first expression unit comprises a first nucleic acid sequence that has the nucleotide sequence of SEQ ID NO:5.
- 185. (Amended) The expression system of claim 181, wherein said second expression unit comprises a second nucleic acid sequence that encodes a polypeptide that comprises a contiguous sequence of at least 16 amino acids from SEQ ID NO:4, of at least 20 amino acids from SEQ ID NO:45, of at least 20 amino acids from SEQ ID NO:47 or of at least 125 amino acids from SEQ ID NO:50.
- 186. The expression system of claim 185, wherein said second expression unit comprises a second nucleic acid sequence that encodes a polypeptide that has the amino acid sequence of SEQ ID NO:4.

- 187. The expression system of claim 185, wherein said second expression unit comprises a second nucleic acid sequence that encodes a polypeptide that has the amino acid sequence of SEQ ID NO:45.
- 188. The expression system of claim 185, wherein said second expression unit comprises a second nucleic acid sequence that encodes a polypeptide that has the amino acid sequence of SEQ ID NO:47.
- 189. The expression system of claim 185, wherein said second expression unit comprises a second nucleic acid sequence that encodes a polypeptide that has the amino acid sequence of SEQ ID NO:50.
- 190. (Amended) The expression system of claim 181, wherein said second expression unit comprises a second nucleic acid sequence that encodes a polypeptide and that comprises the nucleotide sequence of the coding sequence of a cDNA molecule present in a nucleic acid library, wherein the cDNA molecule hybridizes to a probe having the sequence of the complement of SEQ ID NO:3, SEQ ID NO:43 or SEQ ID NO:48 under conditions of high stringency comprising hybridization in 50% formamide, $5\times$ Denhardts' solution, $5\times$ SSC, 25 mM sodium phosphate, 0.1% SDS and 100 µg/ml of denatured salmon sperm DNA at 42°C for 16 h followed by 1h sequential washes with 0.1× SSC, 0.1% SDS solution at 60°C.
- 191. The expression system of claim 190, wherein said second expression unit comprises a second nucleic acid sequence that has the nucleotide sequence of SEQ ID NO:44.
- 192. The expression system of claim 190, wherein said second expression unit comprises a second nucleic acid sequence that has the nucleotide sequence of SEQ ID NO:46.
- 193. The expression system of claim 190, wherein said second expression unit comprises a second nucleic acid sequence that has the nucleotide sequence of SEQ ID NO:49.
- 194. (Amended) The expression system of claim 181, wherein said second expression unit comprises a second nucleic acid sequence that encodes a polypeptide that exhibits between 91% and about 95% identity to the amino acid sequence set forth in SEQ ID NO:4, SEQ ID NO:45, SEQ ID NO:47 or SEQ ID NO:50.

- 195. The expression system of claim 181, wherein said first and said second expression units are comprised in a single expression vector.
- 196. The expression system of claim 181, wherein said first and said second expression units are each comprised in a separate expression vector.
- 197. The expression system of claim 181, wherein said expression system is comprised within a recombinant host cell.
- 198. A recombinant host cell comprising an isolated nucleic acid molecule in accordance with claim 110, claim 113, claim 149 or claim 152.
- 199. The recombinant host cell of claim 198, wherein said cell is a prokaryotic host cell.
- 200. The recombinant host cell of claim 198, wherein said cell is a eukaryotic host cell.
- 201. The recombinant host cell of claim 200, wherein said cell is a mammalian host cell.
- 202. The recombinant host cell of claim 198, wherein said cell further comprises an HIV Tat protein.
- 203. (Amended) The recombinant host cell of claim 198, wherein said cell comprises an isolated nucleic acid molecule comprising a nucleic acid sequence that encodes:
 - (a) a polypeptide having the amino acid sequence of SEQ ID NO:2; or
 - (b) a polypeptide that comprises a contiguous sequence of at least 16 amino acids from SEQ ID NO:4, of at least 20 amino acids from SEQ ID NO:45, of at least 20 amino acids from SEQ ID NO:47 or of at least 125 amino acids from SEQ ID NO:50.

- 204. (Amended) The recombinant host cell of claim 198, wherein said cell comprises an isolated nucleic acid molecule comprising a nucleic acid sequence that encodes a polypeptide that comprises a contiguous sequence of at least 16 amino acids from SEQ ID NO:4, of at least 20 amino acids from SEQ ID NO:47 or of at least 125 amino acids from SEQ ID NO:50.
- 205. (Amended) The recombinant host cell of claim 198, wherein said cell comprises an isolated nucleic acid molecule comprising a nucleic acid sequence that encodes a P-TEFb large subunit protein, wherein said P-TEFb large subunit protein binds to a P-TEFb kinase subunit protein to form a P-TEFb enzyme complex that promotes transcription elongation and wherein said nucleic acid molecule comprises the nucleotide sequence of:

the coding sequence of a cDNA molecule present in a nucleic acid library, wherein the cDNA molecule hybridizes to a probe having the sequence of the complement of SEQ ID NO:3, SEQ ID NO:43 or SEQ ID NO:48 under conditions of high stringency comprising hybridization in 50% formamide, $5\times$ Denhardts' solution, $5\times$ SSC, 25 mM sodium phosphate, 0.1% SDS and 100 μ g/ml of denatured salmon sperm DNA at 42°C for 16 h followed by 1h sequential washes with 0.1× SSC, 0.1% SDS solution at 60°C.

- 206. (Amended) The recombinant host cell of claim 198, wherein said cell comprises an isolated nucleic acid molecule comprising a nucleic acid sequence that encodes a P-TEFb large subunit protein that exhibits at least 90% identity to the amino acid sequence set forth in SEQ ID NO:4, SEQ ID NO:45, SEQ ID NO:47 or SEQ ID NO:50, wherein said P-TEFb large subunit protein binds to a P-TEFb kinase subunit protein to form a P-TEFb enzyme complex that promotes transcription elongation.
- 207. (Amended) The recombinant host cell of claim 198, wherein said cell comprises an isolated nucleic acid molecule comprising:
 - (a) a first nucleic acid sequence that encodes a P-TEFb small subunit protein that has kinase activity and binds to a P-TEFb large subunit protein to form a P-TEFb enzyme complex that promotes transcription elongation, wherein said P-TEFb small subunit protein exhibits at least 90% identity to the amino acid sequence set forth in SEQ ID NO:2 or SEQ ID NO:6; and
 - (b) a second nucleic acid sequence that encodes a P-TEFb large subunit protein that binds to a P-TEFb kinase subunit protein to form a P-TEFb enzyme complex that promotes transcription elongation, wherein said P-TEFb large subunit protein exhibits at least 90% identity to the amino acid sequence set forth in SEQ ID NO:4, SEQ ID NO:45, SEQ ID NO:47 or SEQ ID NO:50.

- 208. A recombinant host cell that comprises an expression system in accordance with claim 157 or claim 181.
- 211. The recombinant host cell of claim 208, wherein said cell is a prokaryotic host cell.
- 212. The recombinant host cell of claim 208, wherein said cell is a eukaryotic host cell.
- 213. The recombinant host cell of claim 212, wherein said cell is a mammalian host cell.
- 214. The recombinant host cell of claim 208, wherein said cell further comprises an HIV Tat protein.
- 215. The recombinant host cell of claim 208, wherein said expression system comprises a first and second expression unit comprised in a single expression vector.
- 216. The recombinant host cell of claim 208, wherein said expression system comprises a first and second expression unit each comprised in a separate expression vector.